OK TO ENTER: /FOH/

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Amendments to the Claims:

This listing of the claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

- 1 (Cancelled).
- 2 (Previously Presented). An isolated protein which is capable of binding to tumor necrosis factor receptorassociated 2 protein (TRAF2), said protein comprising:
 - (A) a polypeptide of SEQ ID NO:3; or
- (B) a variant that has no more than ten amino acid changes from the amino acid sequence of SEQ ID NO:3, wherein said variant is capable of binding to TRAF2.
- 3 (Original). The isolated protein of claim 2, which is a protein comprising the amino acid sequence of SEQ ${
 m TD}$ ${
 m NO}(3)$.
 - 4-19 (Cancelled).
- 20 (Previously Presented). A composition comprising the isolated protein of claim 2 and a pharmaceutically acceptable excipient, diluent, or auxiliary agent.
- 21 (Previously Presented). A molecule having the binding portion of an antibody capable of binding to the portion of said isolated protein of claim 2 that is said polypeptide of (A) or said variant of (B).

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 $% \left(0\right) =0$ (Original). The molecule of claim 21, which is an antibody.

23 (Original). The molecule of claim 22, wherein said antibody is a monoclonal antibody.

24 (Previously Presented). A composition comprising the molecule of claim 21, and a pharmaceutically acceptable excipient, diluent, or auxiliary agent.

25-37 (Cancelled).

38 (Previously Presented). An isolated protein in accordance with claim 2, wherein said protein and said variant are each capable of binding to a component of the NF-xB complex selected from the group consisting of IKappaB kinase complex associated protein (IKAP), IKappaB kinase-alpha (IKK-alpha), IKappaB kinase-beta (IKK-beta), IKappaB kinase-gamma (IKK-gamma) and NF-xB inducing kinase (NIK).

39-41 (Cancelled).

42 (Previously Presented). An isolated protein in accordance with claim 2, comprising a variant of the polypeptide of SEQ ID NO:3, which variant has no more than ten amino acid changes from the amino acid sequence of SEQ ID NO:3, and which variant is capable of binding to TRAF2.

43 (Cancelled).

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- 44 (Previously Presented). A molecule having the binding portion of an antibody capable of binding to the polypeptide of SEQ ID NO:3.
- 45 (Previously Presented). The molecule of claim 44, which is an antibody.
- 46 (Previously Presented). The molecule of claim 45, wherein said antibody is a monoclonal antibody.
 - 47 (Cancelled).
- 48 (Previously Presented). The isolated protein of claim 2, wherein said variant of (B) has no more than five amino acid changes from the amino acid sequence of SEQ ID NO:3.
- 49 (Previously Presented). The isolated protein of claim 2, wherein each said change from the amino acid sequence of SEQ ID NO:3 is a conservative substitution selected from among the substitutions in the following list:

Original

Residue	Substitution
Ala	Gly;Ser
Arg	Lys
Asn	Gln;His
Asp	Glu
Cys	Ser

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Gln Asn Glu Asp Gly Ala; Pro His Asn:Gln Ile Leu; Val Leu Ile; Val Lys Arg; Gln; Glu Met Leu; Tvr; Ile Phe Met;Leu;Tyr Ser Thr Thr Ser Trp Tyr Tvr Trp; Phe

or a conservative substitution that is an exchange within one $% \left(1\right) =\left(1\right) +\left(1$

Ile;Leu

of the following five groups: Small aliphatic, nonpolar or

Val

slightly polar residues: Ala, Ser, Thr, Pro, Gly;

Polar negatively charged

residues and their amides: Asp, Asn, Glu, Gln;

Polar, positively charged residues: His, Arg, Lys;

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Large aliphatic nonpolar residues: Met, Leu, Ile, Val, Cys; and

Large aromatic residues: Phe, Tyr, Trp.

- 50 (Previously Presented). The isolated protein of claim 49, wherein said variant has no more than 5 of said amino acid changes from the amino acid sequence of SEQ ID NO:3.
 - 51 (New). An isolated protein comprising:
 - A) a polypeptide of SEQ ID NO: 3; or
- B) a variant that has no more than 10 amino acid changes from the amino acid sequence of SEO ID NO:3.
- 52 (New). The isolated protein of claim 51, wherein said variant has no more than 5 of said amino acid changes from the amino acid sequence of SEQ ID NO:3.